Fig. 1A

SS 70 10 20 30 40 45 50 15 25 GANTTOGGCA CGACCAAACA GTCATTAGTG GACGACTCTA TTGTAATAAA CTGTGCTTTA AAATGTAAAC 125 130 95 100 105 110 115 120 80 90 CAGGGAGOGT TITTTTTCCT CACATTGTCC AGAAGCAACC TITCTTCCTG AGCCTGGATT AATC ATG 180 185 165 170 175 190 145 150 155 160 AGA GAG CTC GTC AAC ATT OCA CTG TTA CAG ATG CTC ACC CTG GTT GOC TTC AGC GGG r o h r T L λF I P L N 205 215 220 225 230 235 240 245 250 210 ACC GAG AAA CTT CCA AAA GCC CCT GTC ATC ACC ACG CCT CTT GAA ACT GTA GAT GCC T E K L P K A P V I T T P L E T V D A> 295 300 270 275 280 285 290 265 TTA GTT GAA GAA GTG GOG ACT TTC ATG TGC GOC GTG GAA TOC TAC COT CAG CCT GAA Y H CAVE S ₽Q LVEEVATF 320 325 330 335 340 345 350 355 315 ATT TOT TOG ACC AGA AAT AAA ATT CTC ATC AAG CTG TTT GAC ACC CGC TAC AGC ATC IKLFD T R SWTRNKIL **3**85 **3**90 395 400 410 370 375 380 405 415 420 OGA GAG NAC GGT CAG CTC CTC ACC ATC CTG AGT GTG GAG GAC AGT GAT GAT GGC ATC RENGQLLTILS VEDSDDG I> 450 455 460 465 470 430 440 445 435 TAC TGC TGC ACA GOC AAC AAT GGA GTG GGA GGG GTG GAA AGT TGT GGC GCC CTG GGA A E YCCTANNG 495 500 SOS 510 515 520 525 530 485 **4**90 CAA GTG AAG ATG AAG CCT AAA ATA ACT CGT CCT CCC ATC AAT GTA AAA ATA ATT GAG QVKHKPKITRPP I N 585 **S**90 550 SSS 560 **S6**S 570 **575 580** GGA TTG AAA GCA GTC CTA CCG TGC ACT ACG ATG GGT AAC CCC AAG CCA TCC GTG TCC LKAVLPCTTHGNPKP 615 620 625 630 635 640 600 605 645 650 595 610 TGG ATT ANG GGG GAC AGT GCT CTC AGG GAA AAT TOO AGG ATT GCA GTT CTT GAA TCT A L R ENSR I 685 670 675 680 690 695 700 GGG AGT TTA AGG ATC CAT AAT GTG CAA AAG GAA GAC GCA GGA CAG TAC OGA TGT GTG н к v K E D C V> R I Q G Q Y 725 735 745 740 750 755 720 730 GCA AAA AAC AGC CTG GGC ACA GCT TAC TOO AAA CTG GTG AAG CTG GAA GTG GAG GTT SKL T Y ĸ S G 795 800 765 770 775 780 785 790 805 610 TIT OCA AGA ATC CTG OGT OCT OCT GAA TOO CAC AAT GTC ACC TIT GGT TOO TIT GTA ARILRAPE SHHVT F\_G S

Fig. 1B

840 845 850 855 860 825 830 835 865 ACC CTA CGC TGC ACA GCA ATA GGC ATG CCT GTC CCC ACC ATC AGC TGG ATT GAA AAC TLRCTAIGMPVPTISKIE N> 900 905 910 915 920 925 930 935 880 885 890 895 GGA AAT GCT GTT TCT TCA GGT TCC ATT CAA GAG AAT GTG AAA GAC CGA GTG ATT GAC G M A V S S G S I Q E M V K D R V I D> **950** 955 <u>960</u> 965 970 985 975 980 940 TCA AGA CTC CAG CTC TTT ATC ACA AAG CCA GGA CTC TAC ACA TGC ATA GCT ACC AAT SRLQLFITKPGLYTCIATN> 1005 1010 1015 1020 1025 1030 1035 1040 1045 arg cat gga gag aaa ttc agt acc gca aag gct gca gcc act gtc agt ata gca gaa K H G E K F S T A K A A T V S I A E> 1075 1080 1085 1090 1095 1100 1050 1055 1060 1065 1070 tgg agc aaa tca cag aaa gaa agc aaa ggc tac tgt gcc cag tac aga ggg gag gtg W S K S Q K E S K G Y C A Q Y R G E V> 1110 1115 1120 1125 1130 1135 1140 1145 1150 1155 1160 TGT GAT GOO GTC CTG GTG AAA GAC TOT CTT GTC TTC TTC AAC AOC TOO TAT COO GAC C D A V L V K D S L V F F N T S Y P D> 1170 1175 1180 1185 1190 1195 1200 1205 1210 1215 1220 CCT GAG GAG GCC CAA GAG CTG CTG ATC CAC ACT GCG TGG AAT GAA CTC AAG GCT GTG P E E A Q E L L I H T A W N E L K A V>1225 1230 1235 1240 1245 1250 1255 1260 1265 1270 AGC CCA CTC TGC CGA CCA GCT GCC GAG GCT CTG CTG TGT AAT CAC CTC TTC CAG GAG S P L C R P A A E A L L C N H L F Q E> 1315 1320 1325 1330 1285 1290 1295 1300 1305 1310 TGC AGC OCT GGA GTG CTA OCT ACT OCT ATG OOC ATT TGC AGA GAG TAC TGC TTG GCA C S P G V L P T P M P I C R E Y C L A> 1335 1340 1345 1350 1355 1360 1365 1370 1375 1380 1385 1390 GTA AAG GAG CTC TTC TGT GCA AAG GAA TGG CTG GCA ATG GAA GGG AAG ACC CAC GGC V K E L F C A K E W L A M E G K T H R> 1405 1410 1415 1420 1425 1430 1435 1440 1445 1395 1400 GGA CTC TAC AGA TOO GGG ATG CAT TTC CTC COG GTC COG GAG TGC AGC AAG CTT COC GLYRSGHHF PECSK L P> 1465 1470 1475 1480 1485 1490 1495 1500 1505 1455 1460 AGC ATG CAC CAG GAC CCC ACA GCC TGC ACA AGA-CTG COG TAT TTA GAT TAT AAA AAA PTACTRLPYLDY 1510 1515 1520 1525 1530 1535 1540 1545 1550 1555 1560 GAA AAC ATA ACA TTC CCG TCC ATA ACG TCC TCC AAG CCG AGC GTG GAC ATT CCA ENITTEPSITSSKPSVDIP> 1565 1570 1575 1580 1585 1590 1595 1600 1605 1610 1615 AAC CTG OCT OCC TOC ACG TOT TOC TTC GOO GTC TOG CCT GOG TAC TOC ATG ACT GTC TSSFAVSPAYSHT'V> r b v z

Fig. 1C

1620 1625 1630 1645 1650 1655 1660 1635 1640 1665 1670 ATC ATC TOO ATC ATG TOO TGC TIT GOG GTG TIT GCT CTC CTC ACC ATC ACT ACT CTC IISIMSCFAVFALLTITT L> 1680 1685 1690 1695 1700 1705 1710 1715 1720 tat toc toc cga agg agg agg tog aaa aat aag aaa aga gag tog gca gcg gtg Y C C R R R R E W K N K K R E S A A V> 1735 1740 1745 1750 1755 1760 1765 1770 1775 1780 1785 1790 ACC CTC ACC ACA TTG CCT TCC GAG CTC CTG CTG GAC AGG CTG CAT CCC AAC CCC ATG TTLPSELLLDRLHPNPH> 1800 1805 1810 1815 1820 1825 1830 1835 1840 TAC CAG AGG ATG OCA CTC CTT CTG AAT COC AAG TTG CTC AGC CTG GAG TAT COG AGG YQRHPLLLNPKLLSLE 1855 1860 1865 1870 1875 1880 1885 1890 1895 AAT AAC ATC GAG TAT GTC AGA GAC ATC GGA GAG GGA GOG TIT GGA AGG GTC TIT CAA N N I E Y V R D I G E G A F G R V F Q> 1905 1910 1915 1920 1925 1930 1935 1940 1945 1950 1955 1960 gog agg goc oca ggc ttg ctt cot tat gaa coc ttc act atg gtg gct gtg aag atg ARAPGLLPYEPFTMVAVKED 1965 1970 1975 1980 1985 1990 1995 2000 2005 2010 2015 CTG AAG GAG GAC TOC GCA GAT ATG CAG GCA GAC TIT CAG AGG GAG GCA GOC CTC LKEEAS ADMQADFQRE 2020 2025 2030 2035 2040 2045 2050 2055 2060 2065 2070 2075 ATG GOG GAG TIT GAC AAC COC AAC ATT GTG AAG CTC TTA GGT GTG TGT GCT GTT GGG NPNIVKLLGVCA 2085 2090 2095 2100 2105 2110 2115 2120 AAG CCA ATG TGC CTG CTC TTT GAA TAT ATG GCC TAT GGT GAC CTC AAT GAG TTC CTC KPMCLLFEYMAYGDLNEFL> 2135 2140 2145 2150 2155 2160 2165 2170 2175 2180 2185 OGA AGO ATG TOO OCT CAC ACT GTG TGC AGO CTC AGO CAC AGT GAC CTG TOO AGG AGG R S M S P H T V C S L S H S D L S T R> 2190 2195 2200 2205 2210 2215 2220 2225 2230 2235 2240 GCT CGG GTG TCC AGC CCT GGT CCT CCA CCC CTG TCT TGT GCG GAA CAG CTC TGT ATT A R V S S P G P P P L S C A E Q L C I> 2250 2255 2260 2265 2270 2275 2280 2285 2290 GOC AGG CAA GTG GCA GCT GGC ATG GCC TAC CTG TOG GAG CGC AAG TTT GTC CAT CGG A R Q V A A G M A Y L S E R K F V H R> 2310 2315 2320 2325 2330 2335 2340 2345 2350 2355 2360 GAC TTA GCT ACC AGG AAC TGC CTG GTT GGA GAG AAC ATG GTG GTG AAA ATT GCA GAC D L A T R N C L V G E N M V V K I A D> 2370 2375 2380 2385 2390 2395 2400 2405 2410 TIT GGC CTC TCT AGG AAC ATC INC TCC GCA GAC TAC TAC AAA GCT GAT GGA AAC GAT FGLSRNIYSADYYKADGND>

Fig. 1D

2425 2430 2435 2440 2445 2450 2455 2460 2465 2470 GCT ATA OCT ATC OGC TGG ATG OCA OOC GAG TCT ATC TTC TAC AAC OGC TAC AOC ACG AIPIRWEP ESIF Y N R Y T T> 2490 2495 2500 2475 2480 2485 2505 2510 2515 2520 2525 2530 GAG TCA GAT GTG TGG GCT TAT GGC GTG GTC CTC TGG GAG ATC TTC TCC TAT GGA CTG X Y G VVLWE ŜĎ W V I E SYG 2535 2540 2545 2550 2555 2560 2565 2570 2575 CAG CCC TAC TAT GGA ATG GCC CAT GAG GAG GTC ATT TAC TAT GTG AGA GAT GGT AAC H E I Y Y V R D E V Y G K H 2595 2600 2605 2610 2615 2620 2625 2630 2635 2640 2645 ATC CIT GCC TGC CCT GAG AAC TGT CCC TTG GAA CTG TAC AAC CIT ATG CGC CTA TGT ACPENCPLEL YNLHR 2670 2675 2680 2685 2690 2695 2655 2660 2665 TOG AGC AAG CTG CCT GCA GAC AGA CCC AGC TTC TGC AGT ATC CAC CGG ATC CTG CAG W S K L P A D R P S F C S I H R I L Q> 2705 2710 2715 2720 2725 2730 2735 2740 2745 2750 2755 2760 CGC ATG TGC GAG AGA GCA GAG GGA ACG GTA GGC GTC TAA GGTTGACCA TGCTCAAACA TV R A E G G V M C E 2765 2770 2775 2780 2785 2790 2795 2800 2805 2810 2815 2820 2825 2830 ACACCAGGA GGATCTTTTC AGACTGCGAG CTGGAGGGAT CCTAAAGCAG AGGGCGATA AGARCAGATA 2835 2840 2845 2850 2855 2860 2865 GGAAGAGTIT ATCTCAGGCA GCACGINCAG TIGGITGIT

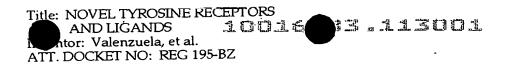
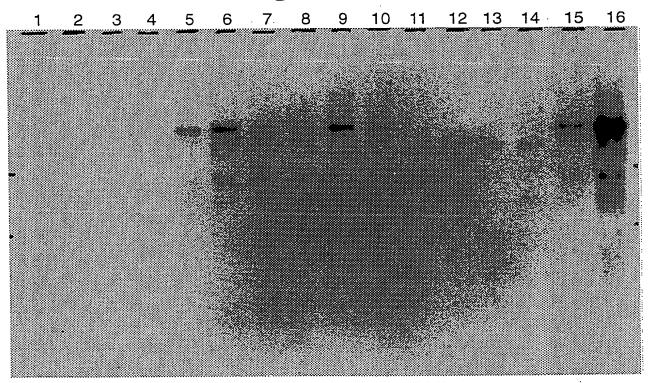


Fig.2.



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77/9

VII. DOCKEL NO: KEC 162-BZ Iuventot: Valenzuela, et al. AND LICANDS Title: NOVEL TYROSINE RECEPTORS. Title: NOVEL TYROSINE RECEPTORS
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ntor: Valenzuela, et al.
ATT. DOCKET NO: REG 195-BZ

7/24

Fig. 4A

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P CCA T	K	ATA I	ACT T.	R	CCT P	ecc g (40	I	N AAA	GTG V 450 CCA	K	I	ATA I	GAG E 60 TGG	G	TTA L	470	<b>A</b> -	V	CTA L 480
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P CCA T	K rct.	ATA I 4: ACT T	ACT T.	R ATG	ect b	CCC P (40 AAT	ccc	N AAA	GTG V 450 CCA	K TCA	I	ATA I TOT	GAG E 60 TGG	G ATA	TTA L AAG K	AAA K 470 GGA	A. GAC	Yec	CTA L 480 CCT
P CCA T P	K C	ATA I 4: ACT T	ACT T.	R ATG H	CCT P	CCC E I40 AAT N	ccc b	AAA K	GTG V 450 CCA P	K TCA S	GTG V	ATA I TCT S	GAG E 50 TGG W	G ATA I	TTA L AAG K	470 GGA G	GAC D	AGC S	CTA L 480 CCT P
CTC A	K C C	ATA I 4: ACT T 4:	ACT T.	R ATG H	CCT P	CCC P (40 AAT N 600	CCC P	N AAA K GTT	GTG V 450 CCA P 510	K TCA S	GTG V	ATA I TCT S SS GGG	GAG E F TGG W	G ATA I	TTA L AAG K	470 GGA G	GAC D	AGC S	CTA L 480 CCT P 540 GTA
CTC A	K C C	ATA I 4: ACT T 4: GAA E	ACT T.	ATG H TCC S	CCT P GCT G	ATT	I CCC P	AAA K GTT V	GTG V 450 CCA P 510 CTT L	K TCA S GAA E	GTG V	ATA I TCT S SS GGG G	GAG E 50 TGG W 20 AGC	G ATA I	TTA L AAG K AGG R	AAA K 470 GGA G 530 ATT I	GAC D CAT H	AGC S	CTA L 480 CCT P 540 GTA V
CTC A	K C C	ATA I 4: ACT T 4: GAA E	ACT T.	R ATG H	CCT P GCT G	ATT	I CCC P	AAA K GTT V	GTG V 450 CCA P 510 CTT L	K TCA S GAA E	GTG V	ATA I TCT S SS GGG	GAG E 50 TGG W 20 AGC	G ATA I	TTA L AAG K AGG R	AAA K 470 GGA G 530 ATT	GAC D CAT H	AGC S	CTA L 480 CCT P 540 GTA
CTC A	K C C	ATA I ACT T 49 GAA E 59	ACT T 30 ACA T 90 AAT N	ATG H TCC S	CCT P GGT G	AAT N GOO ATT I	CCC P	AAA K	GTG V 450 CCA P S10 CTT L	TCA S GAA E	GTG V	ATA I TCT S SS GGG G SS	GAG E 50 TGG W 20 AGC S	G ATA I TTG L	AAG K	AAA K 470 GGA G 530 ATT I	GAC D CAT H	AGC S	CTA L 480 CCT P S40 GTA V
CAA A	K CCT. AGG R	ATA I ACT T 49 GAA E 59	ACT T 30 ACA T 90 AAT N	ATG H TCC S	CCT P GGT G CCA R	AAT N SOO ATT I	CCC P  GCA A	AAA K GTT V	GTG V 450 CCA P S10 CTT L 570	TCA S GAA E	GTG V TCT S	ATA I TCT S SS GGG G SS	GAG GAG TGG W 20 AGC S	G ATA I TTG L	AAG K AGG R	AAA K 470 GGA G 530 ATT I	GAC D CAT H	AGC S	CTA L 480 CCT P 540 GTA V 600 TAT
CAA A	K CCT. AGG R	ATA I ACT T 49 GAA E 59	ACT T 30 ACA T 90 AAT N	ATG H TCC S	CCT P GGT G CCA R	AAT N SOO ATT I	CCC P  GCA A  TAT Y	AAA K GTT V CGA R	GTG V 450 CCA P S10 CTT L S70	TCA S GAA E	GTG V TCT S	ATA I TCT S SS GGG G AAA K	GAG E 50 TGG W 20 AGC S	G ATA I TTG L	AAG K AGG R CTC	AAA K GGA G S30 ATT I S90 GGG G	GAC D CAT H	AGC S	CTA L 480 CCT P 540 GTA V 600 TAT
CAA A	K CCT. AGG R	ATA I ACT T GAA E SS	ACT T 30 ACA T 90 AAT N	ATG H TCC S	CCT P CGT CGA R	AAT N SOO ATT I	CCC P  GCA A  TAT Y	AAA K GTT V CGA R	GTG V 450 CCA P S10 CTT L S70	TCA S GAA E	GTG V TCT S	ATA I TCT S SS GGG G AAA K	GAG GAG TGG W 20 AGC S	G ATA I TTG L	AAG K AGG R CTC	AAA K 470 GGA G 530 ATT I 590	GAC D CAT H	AGC S	CTA L 480 CCT P 540 GTA V 600 TAT
CTC A	K C NGG R	ATA I ACT T AS GAA E SS GAA E	ACT T SO ACA T SO GAT N	ATG H TCC S	CCT P CGT G CGA R	AAT N GOO ATT I CAG	CCC P GCA A TAT	AAA K GTT V CGA R	GTG V 450 CCA P S10 CTT L S70 TGT C	K TCA S GAA E GTG V	GTG V TCT S	ATA I TCT S SS GGG G SAAA K	GAG GAG TGG W 20 AGC S AAC N	G ATA I TTG L AGC S	AAG K AGG R CTC	470 6GA G 530 ATT I 590 GGG G	GAC D CAT H	AGC S AAC N	CTA L 480 CCT P 540 GTA V 600 TAT Y
CCA TO A L CAA A Q	K CCT. CC AGG R	ATA I ACT T AS GAA E SS GAA E GTG	ACT T 30 ACA T 80 AAT N 50 GAT D	ATG H TCC S CCA A	CCT P CCA R CCA CCA CCA CCTC	AAT N GOO ATT I GAA GAA	CCC P GCA A TAT Y	AAA K GTT V CGA R	GTG V 450 CCA P 510 CTT L 630 CTT	K TCA S GAA E GTG V	GTG V TCT S	ATA I TOT S SS GGG G SAAA K AGG	GAG GAG TGG W 20 AGC S AAC N AAC N ATC	G ATA I TTG L AGC S	AAG K AGG R CTC L	AAA K 470 GGA G 530 ATT I 590 GGG G	GAC D CAT H	AGC S AAC N AAC A	CTA L 480 CCT P 540 GTA V 600 TAT Y 660
CCA TO A L CAA A Q	K CCT. CC AGG R	ATA I ACT T AS GAA E SS GAA E GTG	ACT T 30 ACA T 80 AAT N 50 GAT D	ATG H TCC S CCA A	CCT P CCA R CCA CCA CCA CCTC	AAT N GOO ATT I GAA GAA	CCC P GCA A TAT Y	AAA K GTT V CGA R	GTG V 450 CCA P 510 CTT L 630 CTT	K TCA S GAA E GTG V	GTG V TCT S	ATA I TOT S SS GGG G SAAA K AGG	GAG GAG TGG W 20 AGC S AAC N AAC N ATC	G ATA I TTG L AGC S	AAG K AGG R CTC L	470 6GA G 530 ATT I 590 GGG G	GAC D CAT H	AGC S AAC N AAC A	CTA L 480 CCT P 540 GTA V 600 TAT Y 660
CCA TO A L CAA A Q	K CCT. CC AGG R	ATA I ACT T GAA E SS GAA E GTG V	ACT T 30 ACA T 80 AAT N 50 GAT D	ATG H TCC S CCA A	CCT P CCA R CCA CCA CCTC CTC	AAT N GOO ATT I GAA GAA E	CCC P  GCA A  TAT Y	AAA K GTT V CGA R GAG E	GTG V 450 CCA P 510 CTT L 570 GTT C	CA S GAA E GIG V	GTG V TCT S	ATA I TOT S SS GGG G S AAA K 6 AGG R	GAG GAG TGG W 20 AGC S AAC N ATC	G ATA I TTG L AGC S	AAG K AGG R CTC L CGG	GGA G S30 ATT I S90 GGG G G G G A G G A G G G G G G G G G	GAC D CAT H	AGC S AAC N AAC A	CTA L 480 CCT P 540 GTA V 600 TAT Y 660
CTC A CAA A Q TCC A	K CCT. AGG R LAG K	ATA I ACT T GAA E SS GAA E GTG V	ACT TO ACA TO GAT DO GAT DO GAT TO	ATG H TCC S GCA A AAG K	CCT P CCA R CCA CCC CTC L	AAT N 600 ATT I 660 CAG Q GAA E	CCC P  GCA A  TAT Y	AAA K GTT V CGA R GAG E	GTG V 450 CCA P 510 CTT L 570 GTT C 630 GTT V 690	K TCA S GAA E GTG V	GTG V TCT S GCA A	ATA I TCT S SS GGG G SI AAA K 6 AGG R 7	GAG GAG TGG W 20 AGC S AAC N ATC I	G ATA I TTG L AGC S	AAG K AGG R CTC	4 AAA K 470 GGA G 530 ATT I 590 GGG G 710	GAC D  CAT H  ACA T	AGC S AAC A GAA E	CTA L 480 CCT P 540 GTA V 600 TAT Y 660 TCC S 720
CAC A	K CCT. AGG R AGG K AAT	ATA I ACT T AS GAA E SS GAA E GTG V GTC	ACT TO SO ACA TO SO GAT DO ACC	ATG H TCC S AAG K	CCT P CCA R CCA CTC L CCC	40 AAT N 600 ATT I 660 CAG Q 680 TCC	CCC P  GCA A  TAT Y  TIT	AAA K GTT V CGA R GAG E	GTG V 450 CCA P 510 CTT L 570 G30 GTT V 690 ACC	CTC	GTG V TCT S GCA A	ATA I TOT S SS GGG G SAAA K 6 AGG R 7 TGT	GAG GAG TGG W 20 AGC S AAC N ATC I 00 ACA	G ATA I TTG L AGC S CTG L	AAG R CTC L CGG	4 AAA K 470 GGA G 530 ATT I 590 GGG G 710	GAC D CAT H	AGC S AAC A CAA E CCT	CTA L 480 CCT P S40 CTA V 600 TAT Y 660 CTC S 720 CTC

Title: NOVEL TYROSINE RECEPTORS
AND LIGANDS 10016 33-413001

I or: Valenzuela, et al. ATT. DOCKET NO: REG 195-BZ

8/24

## Fig. 4B

FIG	}. '	4B																	
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		103	•			040			1050			100	•			070			1080
TXT Y			P			V OCC			CTA L		QTC	CVC H		γ	TGG K	aat Ņ	GAA E	CIG	K
		10:	90		1:	100		;	1110			11:	20		1	130		:	1140
GTA	GTG	AGC	€ CCY	GTC	TGC	œc	CCY	GCT	GCT.	GλG	CCT	TTG	• TTG	TGT	AAC	cvc	ATC	TTC	CAG
v	V	S	Ð	V	С	R	P	X	አ	E	¥	L	L	С	н	H	I	F	Q
•		11!	50 <b>★</b>		1:	160			1170			118	30 •		1	190			1200
GYC E	LCC	AGT S	CCT P	GGA G	GTA V	GIG V	b ccl	ACT T	ECI.	ATT I	CCC	ATT	TGC	AGA R	GAG E	TAC Y	TGC C	TTG L	Y CCY
		12:	10		13	220			1230			12	10		1	250		;	1260
GTA	ΑλG	•	•			•			•				•			•			GGA
																			G
		12	70 •		1:	280			1290			130	00		1	310			1320
									TCC										ATG H
_	_																		
			•			•			•				•			•			1380
																			CTA L
		13	90		1	400			1410				20		1	430			1440
XXX K									AAG K			CTC	GAC						TCC
	-	•	•	•	- •	-	_	_	••	•	_	•	_	-	•		_	-	_

Fig. 4C

;	J.	. •	,																
		149	50		14	160		:	L470			14	80		1	(90		1	1500
			•			•	~~~		•				•	.~	~	4			•
TCC	S	TCT	S	rcc	F	S	V	S			Y	S	H	T	A	I	I	S	I
•	•		•	•	•	_	•			•	-		••	-			_		
		151	LO		19	20		:	L530			15	(0		1	550		1	1560
• • • •		AGC	*	~		•	~~~	~~~	~~~	.~	1771	1 CT	100	~~	TIT	4000	<b>***</b>	CC1	AGA
K	S	AUC S	F	GCA A	I	F	V		L				T			C	C	R	R
••			•		-	•	•	_	_	_	-								
		15	70		19	80		:	1590			16	00		1	610		1	L620
101		CAA	•		3.25			101	Cla	TC1	GC1	GC A	GTA.	*~	~~	ACC.	ACA	CTG	<b>←</b>
R	Ann K	CAA.	# H	nnn K	W		AAA K	R	E	S	λ	₩.	V	T	L	Ť	T	L	P
		-																	
		16	30		16	540		:	1650			16	60		1	670	•	1	L680
47-41	GAG	CIC	TTA	CTA	GAT	NC!	CTT	CAT	<u>~</u>	AAC	œ	ATG	TAC	CAG	AGG	ATG	cca	CTC	CTT
s	_		L	L	D	R			P		P		Y		R	H	₽	L	L
														•	_				
		16	90		1	700			1710			17	20		1	730			1740
CTG	AAC:	<u></u>	***	TTG	CTC	AGC	CTG	GλG	TAT	CCA.	AGG	AAT	AAC	ATT	GAA	TAT	GTG	AGA	GAC
L			ĸ		L				Y			ĸ		I	E	Y	V		D
		17	50		1	760			1770			17	80		1	790			1800
ATY	CCA	GAG	GGA	ccc	777	GGA	AGG	GTG	777	CAA	GCX	AGG	OCX.	CCA	GGC	TTA	CII	ccc	TAT
		Ε		X		G			F								L	P	Y
					_										_				
		18	10		1	820			1830			18	40		1	850 <del>•</del>			1860
Gλλ	CCT	TTC	ACT	ATG	GTG	GCA	GTA	λλG	ATG	CTC	ልአል	GAA	GAA	ecc	TCG	GCA	GAT	ATG	СУУ
	P			H					ĸ					Y		¥"			Q
																_			
		18	70		1:	880			1890			19	00		1	910			1920
ccc	GAC	444	CAG	AGG	GAG	GCX	GCC	cro	ATG	GCA	Gλλ	TTT	GAC	AAC	CCT	AAC	ATT	GTG	AAG
λ				R		A				A						ĸ	I	v	ĸ
															,				
	•	19	30		1.	940			1950			19	60		1	970			1980
CTA	TTA	GGA	GTG	TGT	GCT	GTC	GGG	AAG	CCA	ATG	TGC	CTG	CTC	TTT	GAA	TAC	ATG	GCC	TAT
	L																		Y
					_									•	_				2010
			90 . <b>4</b>		2	000			2010			20	20		2	030			2040
GGT	GAC			. GYC	TTC	CTC	. ccc	AGC	: ATG	TCC	CCI	CAC	ACC	GTG	100	AGC	CTC	AGT	CYC
G		, L																	н
					_										_				
		20	50		2	060			2070	•		20	080		2	090			2100
λGT	GXC	1770	707	· ATC	. AGG	GCT	CAG	GTC	700	: AGC	: cc	. 000	· ccc	. ccx	. 000	cro	TCC	107	GCT
s									s										A
					_				217				40		_	1160			2160
		21	110		2	120			2130	,		21	40		2	2150			2160
GAC	CAC	cm	TGO	_ AT7	r GCC	. AGC	CYC	GTY	GC)	, GCT	r GG(	TA C	GCT	TAC	: CT	TCA	GAJ	CG7	. ANG
																			K

Fig. 4D

TIT GIT CAC CGA GAT TIA GCC ACC AGG AAC TGC CTG GTG GGC GAG AAC ATG GTG GTG AAA F V H R D L A T R H C L V G E N H ATT GOD GAD TIT GOD OTO TOO AGG AAC ATO TAO TOA GOA GAO TAO TAO AAA GOT AAT GAA D F G L S R H I Y S A D Y Y K AAC GAC GCT ATC CCT ATC CGT TGG ATG CCA CCA GAG TCC ATT TTT TAT AAC CGC TAC ACT AIPIRNKPPESIFYNRYT ACA GAG TOT GAT GTG TOG GCC TAT GOC GTG GTC CTC TOG GAG ATC TTC TOC TAT GGC CTG SDVKXYGVVLKEIFSYGL CAG CCC TAC TAT GGG ATG GCC CAT GAG GAG GTC ATT TAC TAC GTG CGA GAT GGC AAC ATC YYGKAKEEVIYYVRDGKI CTC TCC TCC CCT GAG AAC TCC CCC GTG GAG CTG TAC AAT CTC ATG CCT CTA TGT TCG AGC CPENCPVELYNLKRLCWS ANG CTG OCT GCA GAC AGA CCC AGT TTC ACC AGT ATT CAC CGA ATT CTG GAA CGC ATG TGT PADRPSFTSIHRILERHC GAG AGG GCA GAG GGA ACT GTG AGT GTC TAA ERAEGTVSV

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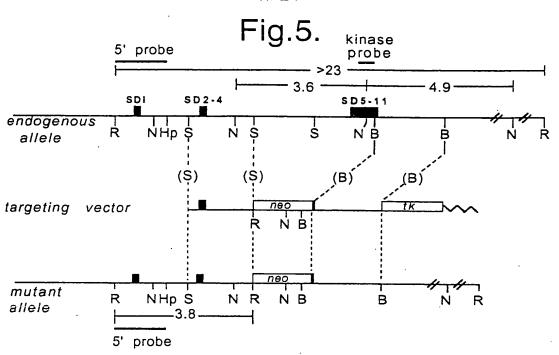
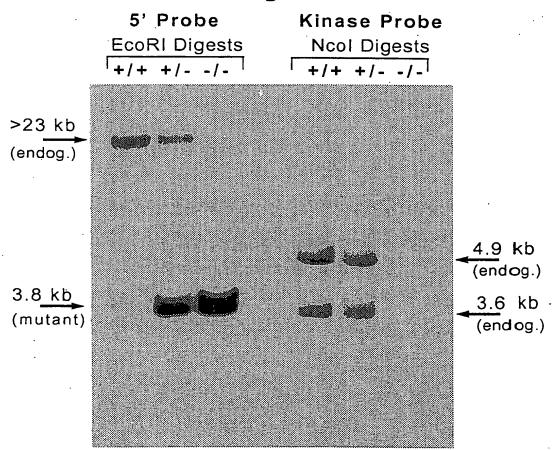
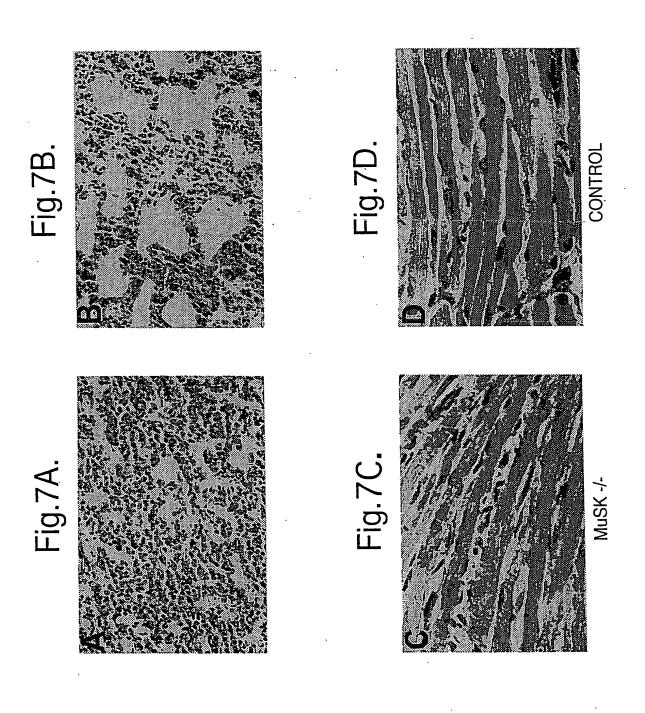
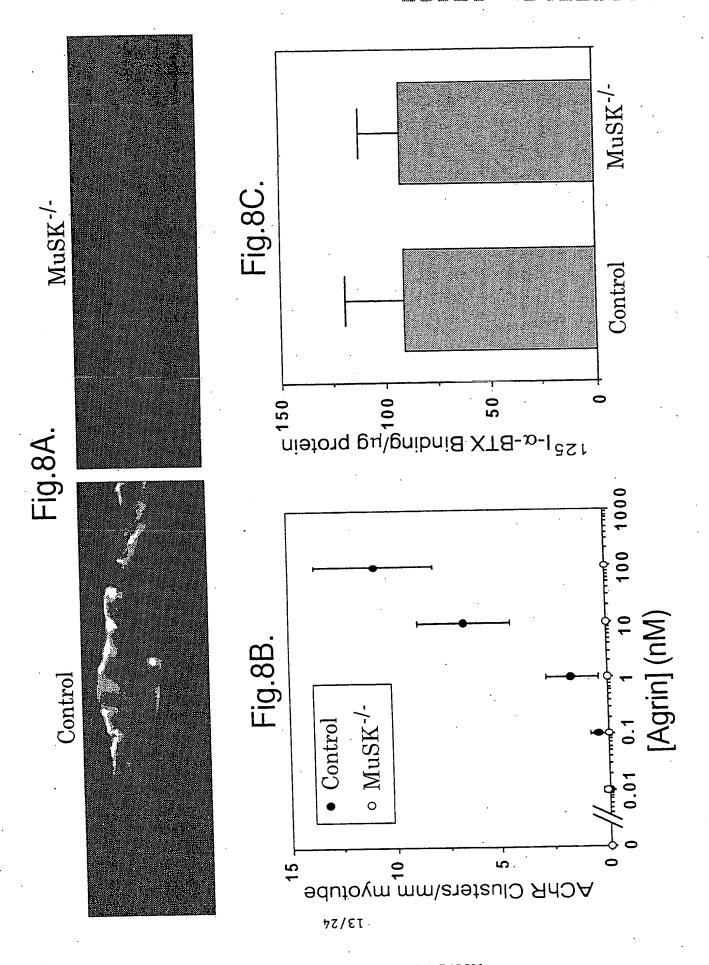


Fig.6.

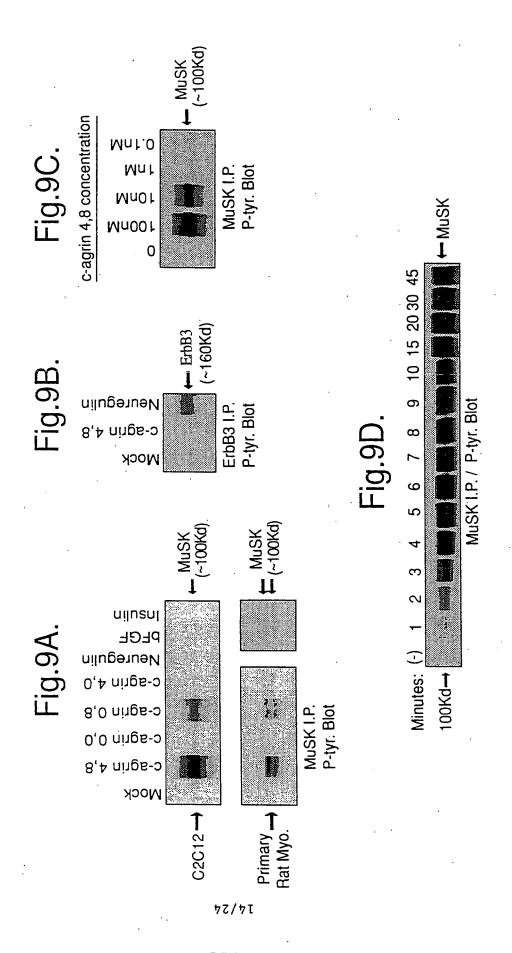




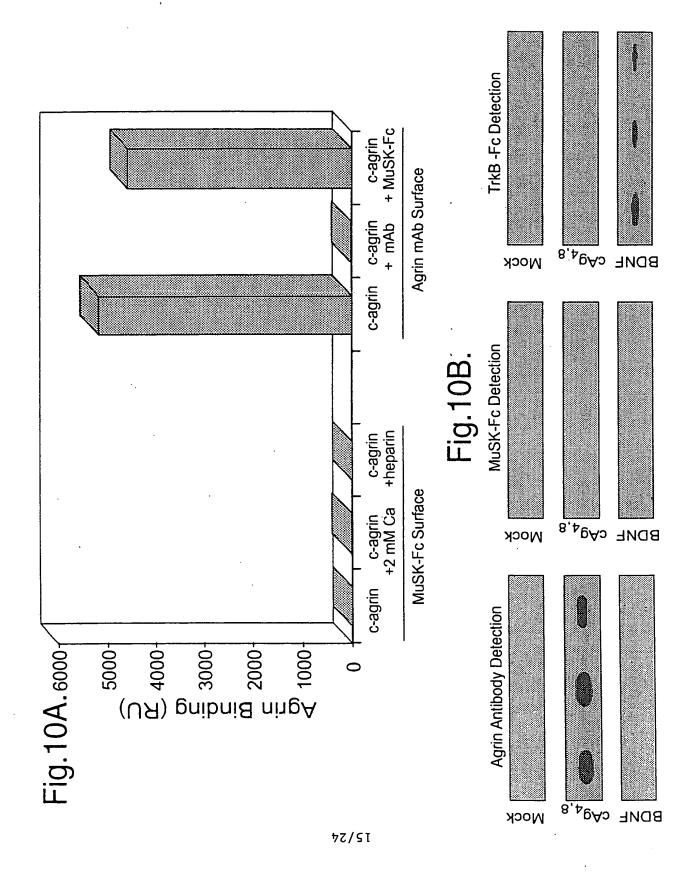
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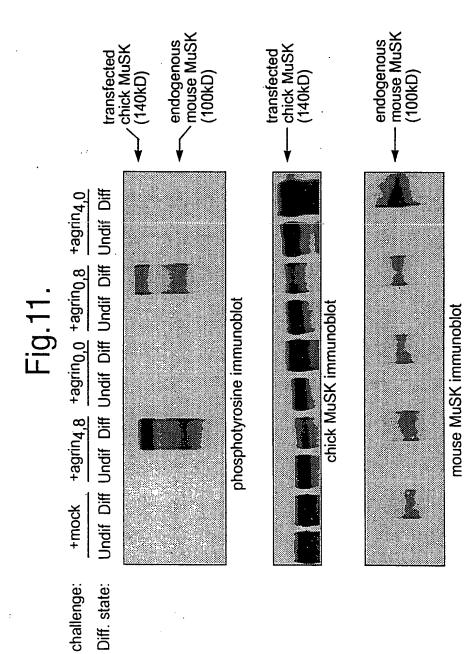
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76/24

0 2 0 4 Z - N 4 F - 0 Z With Soluble B Components B. CNTF Receptor Complex Fc-Tag SYZKGH-O ?Other Receptor Subunits ?Coupling to Extra- & Intra-**Cellular Effector** ?MuSK Dimerization Fig. 12. )SVW β2<sub>)</sub> WNSK β1 C. Formation of Receptor Complex For Agrin A. Formation of CNTF Receptor Complex Associates Binds with MuSk β1 ) SAM Binds omponent Myotube-Associated Binds Receptor Binds

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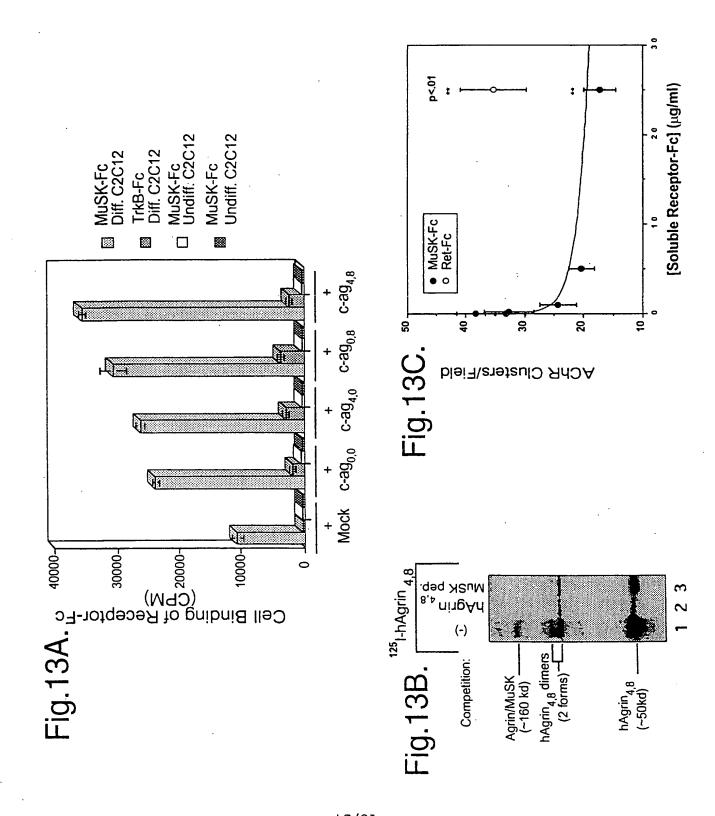


Fig. 14A

1.9	, ,				
10	20	30	40	50	* *
MPPLPLEHRP	RQEPGASMLV	RYFMIPCNIC	LILLATSTIG	FAVLLFLSNY	KPGIHFTPAP
70	80	90	100	110	120
		90			
	•	SVEDPGRASC			
130 * *	140 * *	150°	160 * *	170 * *	180 * *
AQCNQQRRIR					
190	200	210	220	230	240
dsdgvdypse					
					·
* *	* *	270 * *	* *	* *	* *
PENCPAQHTP	ICGDDGVTYE	NDCVMSRIGA	TRGLLLQKVR	SGQCQTRDQC	PETCQFNSVC
310	320	330 * *	340	350	360
		RPVCAQDGHT			
					•
370	380 * *	390 * *	. 400 ★ ★	<b>410</b>	<b>420</b> ★ ★
		AECECQRVCS			
430	440	450 * *	460	470	480
		* * CEVEIGRCVC			
•				,	
490 * *	500 * *	510 * *	520 * *	530 * *	540 * *
THQISLYVAS					GVTYLSACEL
550	560	570	580	590	600 * *
					EDSEDGPCVC
610	620	. 630	640	650	660
* *					* *
DFSCQSVPRS	PVCGSDGVTY	GTECDLKKAR	CESQQELYVA	AQGACRGPTL	APLLPVAFPH
670	680	690	700	710	720 * *
					PGVGGLRCDR
730	. 740	750	760	770	780
					780
		·			KCGQCPDGQV
790	800 * *	810 * *	820 ★ ★	830 * *	840 * *
					VCGSDGVTYG

Fig. 14B

119.17	. D				
850	860 * *	. 870 * *	880 * *	890 * *	900 * *
	RORLDISTOS				
910	920 * *	930	940	950	960
	PLPISPHTTV				
970	980	990	1000	1010	1020
DEELSGDEEA	SGGSGGLEP	PVGSIVVTHG	PPIERASCYN	SPLGCCSDGK	TPSLDSEGSN
1030	1040	1050	1060	1070	1080
	reregaede *				
1090	1100 * *	1110	1120	1130	1140
	KLVRAIVDVH				
1150	1160	1170	1180	1190	1200
	* * * TGAATGTTAA				
	1220 * *				
	RPRTPGHQQP				
	1280 * *				
	HSFLAFPTLR			•	
1330 * *	1340	1350 * *	1360	1370	1380
	PAVLTSLVPV				
1390	1400	1410	1420	1430	1440
					VQSSGVGECG
1450	1460	1470	1480	1490	1500
					AAPCRVLSSG
1510	1520	1530	1540	1550	1560
					EKMALEMVFL
1570	1580	1590	1600	1610	1620
					ALGTWVRVFL
1630	1640	1650	1660	1670	1680
					RGAAVSSGFS

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21/24

# Fig. 14C

1700 1710 GVIQLVSLRG HQLLTQEHVL RAVDVSPFAD HPCTQALGNP CLNGGSCVPR EATYECLCPG GFSGLHCEKG LVEKSVGDLE TLAFDGRTYI EYLNAVIESE KALQSNHFEL SLRTEATQGL ▲Z-site VLWIGKAAER ADYMALAIVD GHLOLSYDLG SOPVVLRSTV KVNTNRWLRI RAHREHREGS LQVGNEAPVT GSSPLGATQL DTDGALWLGG LQKLPVGQAL PKAYGTGFVG CLRDVVVGHR QLHLLEDAVT KPELRPCPTP \*

Fig. 15A ATG TCT GCA CTT CTG ATC CTA GCT CTT GTT GGA GCT GCA GTT GCT GAC A L V L L I L G A A V A -start TAC AAA GAC GAT GAC GAC AAG AAG AGC CCC TGC CAG CCC AAC CCC TGC K K S P ·C D D O P N CAT GGG GCG GCC TGC CGT GTG CTG CCC GAG GGT GGT GCT CAG TGC P С R V L P E G G Α GAG TGC CCC CTG GGG CGT GAG GGC ACC TTC TGC CAG ACA GCC TCG GGG G E G  $\mathbf{T}$ F C P L R Q A CAG GAC GGC TCT GGG CCC TTC CTG GCT GAC TTC AAC GGC TTC TCC CAC F L A D F N G S G P CTG GAG CTG AGA GGC CTG CAC ACC TTT GCA CGG GAC CTG GGG GAG AAG  $\mathbf{T}$ L R G L H F Α R D L G ATG GCG CTG GAG GTC GTG TTC CTG GCA CGA GGC CCC AGC GGC CTC CTG v V F L A R G P S CTC TAC AAC GGG CAG AAG ACG GAC GGC AAG GGG GAC TTC GTG TCG CTG K T D G K G D Y G 0 F GCA CTG CGG GAC CGC CGC CTG GAG TTC CGC TAC GAC CTG GGC AAG GGG E F R R L R Y D L GCA GCG GTC ATC AGG AGC AGG GAG CCA GTC ACC CTG GGA GCC TGG ACC R S R E P V  $\mathbf{T}$ L AGG GTC TCA CTG GAG CGA AAC GGC CGC AAG GGT GCC CTG CGT GTG GGC V S L E R N G R K G A R V G L -Y-insert GAC GGC CCC CGT GTG TTG GGG GAG TCC CCG AAA TCC CGC AAG GTT CCG P R V L G E S P К S K CAC ACC GTC CTC AAC CTG AAG GAG CCG CTC TAC GTA GGG GGC GCT CCC L K E P L Y V G G V L N GAC TTC AGC AAG CTG GCC CGT GCT GCC GTG TCC TCT GGC TTC GAC v K ·L Α R Α Α Α S S GGC GCC ATC CAG CTG GTC TCC CTC GGA GGC CGC CAG CTG CTG ACC CCG Q V S L G G R Q GAG CAC GTG CTG CGG CAG GTG GAC GTC ACG TCC TTT GCA GGT CAC CCC

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TGC ACC CGG GCC TCA GGC CAC CCC TGC CTC AAT GGG GCC TCC TGC GTC

CCG AGG GAG GCT GCC TAT GTG TGC CTG TGT CCC GGG GGA TTC TCA GGA

CCG CAC TGC GAG AAG GGG CTG GTG GAG AAG TCA GCG GGG GAC GTG GAT

EKGLVEKSAG

T

L

С

S

N

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F

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Α

G

23/24

# Fig. 15B

ACC TTG GCC TTT GAC GGG CGG ACC TTT GTC GAG TAC CTC AAC GCT GTG  ${f T}$ F v E Y L F , D G R Α -Z-insert ACC GAG AGC GAA CTG GCC AAT GAG ATC CCC GTC GAG AAG GCA CTG CAG V E K N E I Ρ E Α AGC AAC CAC TTT GAA CTG AGC CTG CGC ACT GAG GCC ACG CAG GGG CTG L R  $\mathbf{T}$ Ė A T Q G E L S H F GTG CTC TGG AGT GGC AAG GCC ACG GAG CGG GCA GAC TAT GTG GCA CTG  ${f T}$ E R Α D Y S G K A GCC ATT GTG GAC GGG CAC CTG CAA CTG AGC TAC AAC CTG GGC TCC CAG Q L S Y G N L G H L CCC GTG GTG CTG CGT TCC ACC GTG CCC GTC AAC ACC AAC CGC TGG TTG V v L R S T V P V N  $\mathbf{T}$ N  $\mathbf{R}$ CGG GTC GTG GCA CAT AGG GAG CAG AGG GAA GGT TCC CTG CAG GTG GGC E Q R E G S L H R Α v AAT GAG GCC CCT GTG ACC GGC TCC TCC CCG CTG GGC GCC ACG CAG CTG V T G S S P L G A Р Α GAC ACT GAT GGA GCC CTG TGG CTT GGG GGC CTG CCG GAG CTG CCC GTG G G L P E WL D G A L GGC CCA GCA CTG CCC AAG GCC TAC GGC ACA GGC TTT GTG GGC TGC TTG P K Α Y G T G F v G L CGG GAC GTG GTG GGC CGG CAC CCG CTG CAC CTG GTG GAG GAC GCC H Ε D A R H P L L L v G GTC ACC AAG CCA GAG CTG CGG CCC TGC CCC ACC CCA TGA V T K P E L R P C P ТP

<sup>24/24</sup> Fig.16.

